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Cytochrome Expression" The contents of which are hereby incorporated by reference--.

Kindly insert the paper copy of the sequence listing provided herewith as pages 40-45 into the specification. Kindly renumber the pages of claims beginning at page 46.

IN THE CLAIMS

Kindly amend the claims as follows:

3. (Amended) A method according to claim 1 comprising screening for said one or more variants in a recognition site for a transcription factor of said regulatory region.

4. (Amended) A method according to claim 1 comprising screening for said one or more variants in an activator protein-3 motif (AP-3) and/or basic transcription element (BTE).

5. (Amended) A method according to claim 1, comprising screening for said one or more variants at at least one of positions -475 or -147 of the transcription regulatory region of the sequence encoding CYP3A5 the sequence of which region is illustrated in Figure 7.

7. (Amended) A method according to claim 1 wherein said DNA is amplified using oligonucleotide molecules which are capable of hybridising selectively to the wild type or variant sequences respectively such that generation of amplified DNA from said respective molecules will indicate whether said wild type or said variant is present.

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